

hybridization with any nucleic acid sequence which will interfere with detecting the presence of the targeted sequence. Probes may include a label capable of detection, where the label is, for example, a radiolabel, fluorescent dye, biotin, enzyme or chemiluminescent compound. Chemiluminescent compounds include acridinium esters which can be used in a hybridization protection assay (HPA) and then detected with a luminometer. Examples of chemiluminescent compounds and methods of labeling probes with chemiluminescent compounds can be found in Arnold *et al.*, U.S. Patent Nos. 4,950,613, 5,185,439 and 5,585,481; and Campbell *et al.*, U.S. Patent No. 4,946,958.

HPA is a detection method based on differential hydrolysis which permits specific detection of the acridinium ester-labeled probe hybridized to the target sequence or amplicon thereof. HPA is described in detail by Arnold *et al.* in U.S. Patent Nos. 5,283,174 and 5,639,604. This detection format permits hybridized probe to be distinguished from non-hybridized probe in solution and includes both a hybridization step and a selection step. In the hybridization step, an excess of acridinium ester-labeled probe is added to the reaction vessel and permitted to anneal to the target sequence or its amplicon. Following the hybridization step, label associated with unhybridized probe is rendered non-chemiluminescent in the selection step by the addition of an alkaline reagent. The alkaline reagent specifically hydrolyzes only that acridinium ester label associated with unhybridized probe, leaving the acridinium ester of the probe:target hybrid intact and detectable. Chemiluminescence from the acridinium ester of the hybridized probe can then be measured using a luminometer and signal is expressed in relative light units (RLU).

After the nucleic acid-based assay is run, and to avoid possible contamination of subsequent amplification reactions, the reaction mixture can be treated with a deactivating reagent which destroys nucleic acids and related amplification products in the reaction vessel. Such reagents can include oxidants, reductants and reactive chemicals which modify the primary chemical structure of a nucleic acid. These reagents operate by rendering nucleic acids inert towards an amplification reaction, whether the nucleic acid is RNA or DNA. Examples of such chemical agents include solutions of sodium hypochlorite (bleach), solutions of potassium permanganate, formic acid, hydrazine, dimethyl sulfate and similar compounds. More details of a deactivation protocol can be found in Dattagupta *et al.*, U.S. Patent No. 5,612,200.

When performed manually, the complexity and sheer number of processing steps associated with a nucleic acid-based assay introduce opportunities for practitioner-error,

exposure to pathogens, and cross-contamination between assays. Following a manual format, the practitioner must safely and conveniently juxtapose the test samples, reagents, waste containers, assay receptacles, pipette tips, aspirator device, dispenser device, and magnetic rack for performing target-capture, while being especially careful not to confuse racks, test samples, assay receptacles, and associated tips, or to knock over any tubes, tips, containers, or instruments. In addition, the practitioner must carefully perform aspirating and dispensing steps with hand-held, non-fixed instruments in a manner requiring precise execution to avoid undesirable contact between assay receptacles, aerosol formation, or aspiration of magnetic particles or other substrates used in a target-capture assay. As a further precaution, the magnetic field in a manually performed target-capture assay is often applied to only one side of the assay receptacle so that fluids can be aspirated through a pipette tip inserted along the opposite side of the assay receptacle. Although applying a magnetic field to only one side of the assay receptacle is a less efficient means for performing a target capture assay, it is designed to prevent magnetic particles from being unnecessarily aspirated as a result of practitioner inaccuracies.

A need exists for an automated diagnostic analyzer which addresses many of the concerns associated with manual approaches to performing nucleic acid-based assays. In particular, significant advantages can be realized by automating the various process steps of a nucleic acid-based assay, including greatly reducing the risk of user-error, pathogen exposure, contamination, and spillage, while significantly increasing through-put volume. Automating the steps of a nucleic acid-based assay will also reduce the amount training required for practitioners and virtually eliminate sources of physical injury attributable to high-volume manual applications.

SUMMARY OF THE INVENTION

The above-described needs are addressed by an automated clinical analyzer constructed and operated in accordance with aspects of the present invention. In general, the automated clinical analyzer integrates and coordinates the operation of various automated stations, or modules, involved in performing one or more assays on a plurality of reaction mixtures contained in reaction receptacles. The analyzer is preferably a self-contained, stand alone unit. Assay specimen materials and reaction receptacles, as well as the various solutions, reagents, and other materials used in performing the assays are preferably stored within the analyzer, as are the waste products generated when assays are performed.

The analyzer includes a computer controller which runs analyzer-controlling and assay-scheduling software to coordinate operation of the stations of the analyzer and movement of each reaction receptacle through the analyzer.

Reaction receptacles can be loaded in an input queue which sequentially presents each receptacle at a pick-up position to be retrieved by a transport mechanism, which automatically transports the reaction receptacles between the stations of the analyzer.

Specimen containers are carried on a first ring assembly, and disposable pipette tips are carried on a second ring assembly. Containers of target capture reagent, including a suspension of solid support material, are carried on an inner rotatable assembly constructed and arranged to selectively agitate the containers or present the containers for access by the probe of an automatic robotic pipette system. Reaction mixtures, including fluid specimen material and target capture reagent, are prepared by the pipette system within each reaction receptacle.

The analyzer further includes receptacle mixers for mixing the contents of a receptacle placed therein. The mixer may be in fluid communication with fluid containers and may include dispensers for dispensing one or more fluids into the receptacle. One or more incubators carry multiple receptacles in a temperature-controlled chamber and permit individual receptacles to be automatically placed into and removed from the chamber. Magnetic separation wash stations automatically perform a magnetic separation wash procedure on the contents of a receptacle placed in the station.

In the preferred method of operation, assay results may be ascertained by the amount of light emitted from a receptacle at the conclusion of the appropriate preparation steps. Accordingly, the analyzer includes a luminometer for detecting and/or quantifying the amount of light emitted by the contents of the reaction receptacle. A deactivation queue may be provided to deactivate the contents of a reaction receptacle placed therein at the conclusion of the assay.

Reaction receptacles can be independently transported between stations by the transport mechanism, and the stations can be operated in parallel to perform different assay procedures simultaneously on different reaction receptacles, thereby facilitating efficient, high through-put operation of the analyzer. Moreover, the present invention facilitates arranging the various stations associated with a nucleic acid-based assay onto a single, contained platform, thereby achieving efficient space utilization.

Other objects, features, and characteristics of the present invention, including the methods of operation and the function and interrelation of the elements of structure, will become